

***Amendments to the Drawings***

The attached sheets of drawings include changes to Figure 2. These sheets, which include Figures 2A-2D, replace the original sheets including Figures 2A-2D. In Figure 2B, nucleotide residues 361-420 and their corresponding amino acid residues have been added.

Attachment: Replacement Sheets

Annotated Sheets Showing Changes

***Remarks***

Figure 2B has been amended to include nucleotide residues 361-420 and their corresponding amino acid residues. Support for this amendment can be found in the original specification at pages 60-63 (SEQ ID NO:3 and SEQ ID NO:4) and original Figure 2A. Accordingly, these changes are believed to introduce no new matter, and their entry is respectfully requested.

***Conclusion***

Prompt and favorable consideration of this Preliminary Amendment is respectfully requested. Applicants believe the present application is in condition for allowance. If the Examiner believes, for any reason, that personal communication will expedite prosecution of this application, the Examiner is invited to telephone the undersigned at the number provided.

Respectfully submitted,

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nucleotide residues  
361-420 and corresponding  
amino acid residues  
added

361 CAATAGTGGGGCTGCAGTGCATGAACACCAGCAGCTACCTCAGCAAGACGTTATT  
N S E G L Q C M N T S T S Y L S K T L F  
421 TGA AATTACAGTGCCTCTCTCTCAAGGCCCAACACAGTAACAATCAGTTTGGCCAATCA  
E I T V P L S Q G P K P V T I S F A N H  
481 CACTTCCTGCCGATGCTCTAACTGGATGTTTACAGACAAGTTCATTCCATTATTAG  
T S C R C M S K L D V Y R Q V H S I I R  
541 ACGTTCCCTGCCAGCAACACTACCACAGTGTGAGGCGGAGCAAGACCTGCCCCACCAA  
R S L P A T L P Q C Q A A N K T C P T N  
601 TTACATGTGGAATAATCACATCTGCAGATGCCCTGGCTCAGGAAGATTTATGTTTCCTC  
Y M W N N H I C R C L A Q E D F M F S S  
661 GGATGCTGGAGATGACTCAACAGATGGATTCCATGACATCTGTGGACCAACAGGAGCT  
D A G D D S T D G G F H D I C G P N K E L

FIG. 2B

721 GGATGAAGACCTGTCACTGTCTGTCAGAGCGGGCTTCGGCCTGCCAGCTGTGGACC  
-----+-----+-----+-----+-----+-----+-----+  
D E E T C C Q C V C R A G L R P A S C G P  
781 CCACAAAGAACTAGACAGAACTCATGCCAGTGTGTCTGTGTAATAAACAACCTCTTCCCCAG  
-----+-----+-----+-----+-----+-----+-----+  
H K E L D R N S C C Q C V C K N K L F P S  
841 CCAATGTGGGCCAACCGAGAATTGATGAAACACACATGCCAGTGTGTATGTAAAGAAC  
-----+-----+-----+-----+-----+-----+-----+  
Q C G A N R E F D E N T C C Q C V C K R T  
901 CTGCCCCAGAAATCAACCCCTAAATCCTGGAAATGTGCCCTGTGAATGTACAGAAAGTCC  
-----+-----+-----+-----+-----+-----+-----+  
C P R N Q P L N P G K C A C E C T E S P  
961 ACAGAAATGCTTGTAAAGGAAGAAGTTCCACCACCAACATGCAGCTGTTACAGACG  
-----+-----+-----+-----+-----+-----+-----+  
Q K C L L K G K K F H H Q T C S C Y R R  
1021 GCCATGTACGAACCGCAGAGGCTGTGAGCCAGGATTTTCATATAGTGAAGAAGTGTG  
-----+-----+-----+-----+-----+-----+-----+  
P C T N R Q K A C E P G F S Y S E E V C

FIG. 2C

